

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/588, 104  
Source: IFWP  
Date Processed by STIC: 08/10/2006

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IFWP

## RAW SEQUENCE LISTING

DATE: 08/10/2006

PATENT APPLICATION: US/10/588,104

TIME: 11:02:40

Input Set : A:\004974.01207 SEQUENCE LISTING.txt

Output Set: N:\CRF4\08102006\J588104.raw

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3 <110> APPLICANT: Bayer HealthCare AG
5 <120> TITLE OF INVENTION: Diagnostics and Therapeutics for Diseases Associated with
6   Puromycin Sensitive Aminopeptidase NPEPPS (NPEPPS)
8 <130> FILE REFERENCE: 004974.01207
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/588,104
C--> 10 <141> CURRENT FILING DATE: 2006-07-31
10 <160> NUMBER OF SEQ ID NOS: 5
12 <170> SOFTWARE: PatentIn version 3.2
14 <210> SEQ ID NO: 1
15 <211> LENGTH: 4177
16 <212> TYPE: DNA
17 <213> ORGANISM: Homo sapiens
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22 ccgcctcctc cctcctcctt tctcgtcttc agccgctcct ctgcgcgcgc cctccacagc 180
23 ctgggcctcg ccgcgatgcc ggagaagagg cccttcgagc ggctgcctgc cgatgtctcc 240
24 cccatcaact acagcctttg cctcaagccc gacttgcctg acttcacctt cgagggcaag 300
25 ctggaggccg ccgcccaggg gaggcaggcg actaatcaga ttgtgatgaa ttgtgctgat 360
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53 atggaggctt ttgtgaatga gcccaattat actgtatgga gcgacctgag ctgtaacctg 2040
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91 &lt;210&gt; SEQ ID NO: 2

92 &lt;211&gt; LENGTH: 919

93 &lt;212&gt; TYPE: PRT

94 &lt;213&gt; ORGANISM: Homo sapiens

96 &lt;400&gt; SEQUENCE: 2

97 Met Trp Leu Ala Ala Ala Pro Ser Leu Ala Arg Arg Leu Leu Phe

98 1 5 10 15

99 Leu Gly Pro Pro Pro Pro Pro Leu Leu Leu Leu Val Phe Ser Arg Ser

100 20 25 30

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101 Ser Arg Arg Arg Leu His Ser Leu Gly Leu Ala Ala Met Pro Glu Lys
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103 Arg Pro Phe Glu Arg Leu Pro Ala Asp Val Ser Pro Ile Asn Tyr Ser
104      50      55      60
105 Leu Cys Leu Lys Pro Asp Leu Leu Asp Phe Thr Phe Glu Gly Lys Leu
106 65      70      75      80
107 Glu Ala Ala Ala Gln Val Arg Gln Ala Thr Asn Gln Ile Val Met Asn
108      85      90      95
109 Cys Ala Asp Ile Asp Ile Ile Thr Ala Ser Tyr Ala Pro Glu Gly Asp
110      100      105      110
111 Glu Glu Ile His Ala Thr Gly Phe Asn Tyr Gln Asn Glu Asp Glu Lys
112      115      120      125
113 Val Thr Leu Ser Phe Pro Ser Thr Leu Gln Thr Gly Thr Gly Thr Leu
114      130      135      140
115 Lys Ile Asp Phe Val Gly Glu Leu Asn Asp Lys Met Lys Gly Phe Tyr
116 145      150      155      160
117 Arg Ser Lys Tyr Thr Thr Pro Ser Gly Glu Val Arg Tyr Ala Ala Val
118      165      170      175
119 Thr Gln Phe Glu Ala Thr Asp Pro Arg Arg Ala Phe Pro Cys Trp Asp
120      180      185      190
121 Glu Pro Ala Ile Lys Ala Thr Phe Asp Ile Ser Leu Val Val Pro Lys
122      195      200      205
123 Asp Arg Val Ala Leu Ser Asn Met Asn Val Ile Asp Arg Lys Pro Tyr
124      210      215      220
125 Pro Asp Asp Glu Asn Leu Val Glu Val Lys Phe Ala Arg Thr Pro Val
126 225      230      235      240
127 Met Ser Thr Tyr Leu Val Ala Phe Val Val Gly Glu Tyr Asp Phe Val
128      245      250      255
129 Glu Thr Arg Ser Lys Asp Gly Val Cys Val Arg Val Tyr Thr Pro Val
130      260      265      270
131 Gly Lys Ala Glu Gln Gly Lys Phe Ala Leu Glu Val Ala Ala Lys Thr
132      275      280      285
133 Leu Pro Phe Tyr Lys Asp Tyr Phe Asn Val Pro Tyr Pro Leu Pro Lys
134      290      295      300
135 Ile Asp Leu Ile Ala Ile Ala Asp Phe Ala Ala Gly Ala Met Glu Asn
136 305      310      315      320
137 Trp Gly Leu Val Thr Tyr Arg Glu Thr Ala Leu Leu Ile Asp Pro Lys
138      325      330      335
139 Asn Ser Cys Ser Ser Ser Arg Gln Trp Val Ala Leu Val Val Gly His
140      340      345      350
141 Glu Leu Ala His Gln Trp Phe Gly Asn Leu Val Thr Met Glu Trp Trp
142      355      360      365
143 Thr His Leu Trp Leu Asn Glu Gly Phe Ala Ser Trp Ile Glu Tyr Leu
144      370      375      380
145 Cys Val Asp His Cys Phe Pro Glu Tyr Asp Ile Trp Thr Gln Phe Val
146 385      390      395      400
147 Ser Ala Asp Tyr Thr Arg Ala Gln Glu Leu Asp Ala Leu Asp Asn Ser
148      405      410      415
149 His Pro Ile Glu Val Ser Val Gly His Pro Ser Glu Val Asp Glu Ile

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150          420          425          430
151 Phe Asp Ala Ile Ser Tyr Ser Lys Gly Ala Ser Val Ile Arg Met Leu
152          435          440          445
153 His Asp Tyr Ile Gly Asp Lys Asp Phe Lys Lys Gly Met Asn Met Tyr
154          450          455          460
155 Leu Thr Lys Phe Gln Gln Lys Asn Ala Ala Thr Glu Asp Leu Trp Glu
156 465          470          475          480
157 Ser Leu Glu Asn Ala Ser Gly Lys Pro Ile Ala Ala Val Met Asn Thr
158          485          490          495
159 Trp Thr Lys Gln Met Gly Phe Pro Leu Ile Tyr Val Glu Ala Glu Gln
160          500          505          510
161 Val Glu Asp Asp Arg Leu Leu Arg Leu Ser Gln Lys Lys Phe Cys Ala
162          515          520          525
163 Gly Gly Ser Tyr Val Gly Glu Asp Cys Pro Gln Trp Met Val Pro Ile
164          530          535          540
165 Thr Ile Ser Thr Ser Glu Asp Pro Asn Gln Ala Lys Leu Lys Ile Leu
166 545          550          555          560
167 Met Asp Lys Pro Glu Met Asn Val Val Leu Lys Asn Val Lys Pro Asp
168          565          570          575
169 Gln Trp Val Lys Leu Asn Leu Gly Thr Val Gly Phe Tyr Arg Thr Gln
170          580          585          590
171 Tyr Ser Ser Ala Met Leu Glu Ser Leu Leu Pro Gly Ile Arg Asp Leu
172          595          600          605
173 Ser Leu Pro Pro Val Asp Arg Leu Gly Leu Gln Asn Asp Leu Phe Ser
174          610          615          620
175 Leu Ala Arg Ala Gly Ile Ile Ser Thr Val Glu Val Leu Lys Val Met
176 625          630          635          640
177 Glu Ala Phe Val Asn Glu Pro Asn Tyr Thr Val Trp Ser Asp Leu Ser
178          645          650          655
179 Cys Asn Leu Gly Ile Leu Ser Thr Leu Leu Ser His Thr Asp Phe Tyr
180          660          665          670
181 Glu Glu Ile Gln Glu Phe Val Lys Asp Val Phe Ser Pro Ile Gly Glu
182          675          680          685
183 Arg Leu Gly Trp Asp Pro Lys Pro Gly Glu Gly His Leu Asp Ala Leu
184          690          695          700
185 Leu Arg Gly Leu Val Leu Gly Lys Leu Gly Lys Ala Gly His Lys Ala
186 705          710          715          720
187 Thr Leu Glu Glu Ala Arg Arg Arg Phe Lys Asp His Val Glu Gly Lys
188          725          730          735
189 Gln Ile Leu Ser Ala Asp Leu Arg Ser Pro Val Tyr Leu Thr Val Leu
190          740          745          750
191 Lys His Gly Asp Gly Thr Thr Leu Asp Ile Met Leu Lys Leu His Lys
192          755          760          765
193 Gln Ala Asp Met Gln Glu Glu Lys Asn Arg Ile Glu Arg Val Leu Gly
194          770          775          780
195 Ala Thr Leu Leu Pro Asp Leu Ile Gln Lys Val Leu Thr Phe Ala Leu
196 785          790          795          800
197 Ser Glu Glu Val Arg Pro Gln Asp Thr Val Ser Val Ile Gly Gly Val
198          805          810          815

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199 Ala Gly Gly Ser Lys His Gly Arg Lys Ala Ala Trp Lys Phe Ile Lys
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201 Asp Asn Trp Glu Glu Leu Tyr Asn Arg Tyr Gln Gly Gly Phe Leu Ile
202      835      840      845
203 Ser Arg Leu Ile Lys Leu Ser Val Glu Gly Phe Ala Val Asp Lys Met
204      850      855      860
205 Ala Gly Glu Val Lys Ala Phe Phe Glu Ser His Pro Ala Pro Ser Ala
206 865      870      875      880
207 Glu Arg Thr Ile Gln Cys Cys Glu Asn Ile Leu Leu Asn Ala Ala
208      885      890      895
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210      900      905      910
211 Lys Ala Ser Pro Pro Thr Val
212      915
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215 <211> LENGTH: 18
216 <212> TYPE: DNA
217 <213> ORGANISM: artificial sequence
219 <220> FEATURE:
220 <223> OTHER INFORMATION: forward primer
222 <400> SEQUENCE: 3
223 tttctctgcc ccctgtgg                                     18
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226 <211> LENGTH: 23
227 <212> TYPE: DNA
228 <213> ORGANISM: artificial sequence
230 <220> FEATURE:
231 <223> OTHER INFORMATION: reverse primer
233 <400> SEQUENCE: 4
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236 <210> SEQ ID NO: 5
237 <211> LENGTH: 33
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239 <213> ORGANISM: artificial sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: probe
244 <400> SEQUENCE: 5
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VERIFICATION SUMMARY

DATE: 08/10/2006

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TIME: 11:02:41

Input Set : A:\004974.01207 SEQUENCE LISTING.txt

Output Set: N:\CRF4\08102006\J588104.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No  
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date